

SEQUENCE LISTING

<110> Remacle, Jose
 Hamels, Sandrine
 Zammattéo, Nathalie
 Lockman, Laurence
 Dufour, Sophie
 Alexandre, Isabelle
 De Longueville, Francoise

<120> IDENTIFICATION OF A LARGE NUMBER OF
 BIOLOGICAL (MICRO)ORGANISMS GROUPS AT DIFFERENT
 LEVELS BY THEIR DETECTION ON A SAME ARRAY

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<150> EP 00870055.1

<151> 2000-03-24

<150> EP 00870204.5

<151> 2000-03-24

<150> US 09/817,014

<151> 2001-03-23

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1B, 1C, 1D, 1E, 2A, 2B, 2C, 4, 6, and 7 sense

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Forward

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<400> 229

agcgaagaat cgggtaaggg t

21

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<223> Chicken capture nucleotide sequence

<400> 230

ccttaacgac tcttatocaa acactatgcc accggggag

39

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<223> Duck capture nucleotide sequence

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ccctaacgac tcttatocaa acactactgc catcggggag

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ccttaacgaa ctctaag

17

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<223> Pig capture nucleotide sequence

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aaagaggagt agaatacacga ttaag

25

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<223> Rabbit capture nucleotide sequence

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<223> Turkey capture nucleotide sequence

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cctaacgact attctccaac cactactgac aacgaggag 39

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<211> 96
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 cytochrome b

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<220>
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<400> 240
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<210> 241
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Sequence

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tcactcggca tgata 15

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gcatggggta tataa 15

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ataagcgtgg acatta 16

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ccttaaatac ctacc 15

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<223> Kenana capture nucleotide sequence

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tgctatagaa gtcac 15

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gaagcatgca taccatctct agca	24
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 agagaagttg aattgactca agga 24

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 gaagcatgtg aaacatctca gtaa 24

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<211> 21
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 actatthcta gccatvcayt a 21

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 aggtaggagc cataaagacc tcg 23

 <210> 270
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 <220>
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 aaggcttaat cagtcggcat caaatgta 28

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 <223> G. macrocephalus capture nucleotide sequence

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28

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aaagcttaatacagtcggcat taaatgta

28

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29

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aaagcctaata cagtcggcctt taattgca 28

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aacgcctact cagtaggctt caaatgca 28

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aaacattcac gctaacggag catctttctt ctttatctgt 40

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<400> 291
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<220>
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<400> 292
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<223> P450-2 consensus primer

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gaaaggggcg tcttggg 17

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gaaaggggcg tcttggg 17

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gctaactgag cacagga 17

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16

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type

<400> 299
ctcggtcacc ccctgc

16

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(deletion)

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15

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16

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mutation

<400> 302
aattatttcc caggaa 16

<210> 303
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type

<400> 303
agcaccctt gaatcc 16

<210> 304
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<220>
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mutation

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agcaccctt gaatcc 16

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<220>
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<400> 305
gaatttgtt aaatggaa 18

<210> 306
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<220>
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<400> 306
gtagtacgga artagaa 17

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saprophyticus)

<400> 307
ggtggtgaaa tggtcc 16

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cgacctgctg tccagct 17

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<220>
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<400> 309
cttcaggacg tatcgacc 18

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<220>
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ttattagact acgctgaag 19

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<220>
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ccatttccgg ccgcgg 16

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<220>
<223> H. influenzae capture probe

<400> 313
gagtttagcaa accacttag 19

<210> 314
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<220>
<223> E. coli capture probe

<400> 314
aactggctgg cttcctg 17

<210> 315
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gtatcaaaga agaaactcaa a 21

<210> 316

ggtatcaaag aaacttctaa a

21

<210> 321

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<212> DNA

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<223> S. saprophyticus capture probe

<400> 321

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20

Sequence